

420



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                   1
                                                                         99
 ttt tgg att tca gcc tcc aga ggt gat att gtg cta act cag tct cca
 Phe Trp Ile Ser Ala Ser Arg Gly Asp Ile Val Leu Thr Gln Ser Pro
                              20
          15
 gcc acc ctg tct gtg act cca gga gat agc gtc agt ctt tcc tgc agg
                                                                         147
 Ala Thr Leu Ser Val Thr Pro Gly Asp Ser Val Ser Leu Ser Cys Arg
                           35
      30
  gcc agc caa att att agc aac aac cta cac tgg tat caa caa aaa tca
                                                                         195
  Ala Ser Gln Ile Ile Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Ser
                       50
  45
  cat gag tot coa agg oft of atc aag tat got too cag too atc tot
                                                                         243
  His Glu Ser Pro Arg Leu Leu Ile Lys Tyr Ala Ser Gln Ser Ile Ser
                                                                         291
  ggg atc ccc tcc agg ttc agt ggc agt gga tca ggg aca gat ttc act
  Gly Ile Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr
                                   85
               80
  ctc agt atc aac agt gtg gag act gaa gat ttt gga atg tat ttc tgt
                                                                          339
   Leu Ser Ile Asn Ser Val Glu Thr Glu Asp Phe Gly Met Tyr Phe Cys
                                100
           95
                                                                          387
   caa cag agt aac agc tgg cct ctc acg ttc ggc tcg ggg aca aag ctg
   Gln Gln Ser Asn Ser Trp Pro Leu Thr Phe Gly Ser Gly Thr Lys Leu
                            115
       110
                                                                          439
   gag atc aaa cgg cgtaagtgtg tcagggtttc acaagaggga ctaaagacat
   Glu Ile Lys Arg
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483

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Val Thr Pro Gly Asp Ser Val Ser Leu Ser Cys Arg Ala Ser Gln Ile 35 40 45

Ile Ser Asn Asn Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro 50 55 60

Arg Leu Leu Ile Lys Tyr Ala Ser Gln Ser Ile Ser Gly Ile Pro Ser 65 70 75 80

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn 85

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 acggtcaccg tctccagt
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  ctt gtt tta aaa ggt gtc cag tgt gaa gtg aag gtg gtg gag tct ggg
                                                                100
  Leu Val Leu Lys Gly Val Gln Cys Glu Val Lys Val Val Glu Ser Gly
  gga ggc tta gtg aag cct gga gcg tct ctg aaa ctc tcc tgt gca gcc
                                                                 148
  Gly Gly Leu Val Lys Pro Gly Ala Ser Leu Lys Leu Ser Cys Ala Ala
                            35
          30
                                                                 196
   tct gga ttc act ttc agt aac tat ggc atg tct tgg gtt cgc cag act
   Ser Gly Phe Thr Phe Ser Asn Tyr Gly Met Ser Trp Val Arg Gln Thr
                        50
      45
   tca gac aag agg ctg gag tgg gtc gca tcc att agt agt ggt ggt gat
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   Ser Asp Lys Arg Leu Glu Trp Val Ala Ser Ile Ser Ser Gly Gly Asp
                                       70
                     65
   60
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agc acc ttc tat gca gac aat gta aag ggc cga ttc acc atc tcc aga Ser Thr Phe Tyr Ala Asp Asn Val Lys Gly Arg Phe Thr Ile Ser Arg 80 85	292
gag aat gcc aag aac acc ctg tac ctg caa atg agt agt ctg aag tct Glu Asn Ala Lys Asn Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser 95	340
gag gac acg gcc ttg tat tac tgt gca aga gac gat cta ttt aac tgg Glu Asp Thr Ala Leu Tyr Tyr Cys Ala Arg Asp Asp Leu Phe Asn Trp 110	388
ggc caa ggc acc act ctc aca gtc tca tca gccaaaacaa cagccccatc Gly Gln Gly Thr Thr Leu Thr Val Ser Ser 125	438
ggtctatcca ctggcccctg tgtgtggaga tacaattggc tcctcggtga ctttaggatg	498
cctggtcaag ggttatttcc ttgagccagt gaccttgacc tggaactctg gatccctgtc	558
cagtggtgtg cacatettee cagetgtett geagtetgae etetacacee teageagete	618
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Ser Asn Tyr Gly Met Ser Trp Val Arg Gln Thr Ser Asp Lys Arg Leu	

50		55	60	)	
Glu Trp Val A 65	Ala Ser Ile 70	Ser Ser Gly	Gly Asp Se	er Thr Phe Tyr	Ala 80
Asp Asn Val I	Lys Gly Arg 85	Phe Thr Ile	Ser Arg Gl 90	lu Asn Ala Lys 95	Asn
Thr Leu Tyr l	Leu Gln Met 100	Ser Ser Leu 105	Lys Ser Gl	lu Asp Thr Ala 110	Leu
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Leu Thr Val 130	Ser Ser				
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tgg att cag Trp Ile Gln 15	gaa acc aad Glu Thr Asi	c ggt gat gt n Gly Asp Va 20	tt gta atg al Val Met	acc cag act cc Thr Gln Thr Pr 25	a ctc 98 o Leu
act ttg tcg Thr Leu Ser 30	g gtt acc at c Val Thr Il	t gga caa co e Gly Gln P: 35	ca gcc tct ro Ala Ser	atc tct tgc aa Ile Ser Cys Ly 40	g tca 146 s Ser
agt cag ago	c ctc tta ta	t agt aat g	ga aaa acc	tat ttg aat tg	g tta 194

Ser Gln Ser Leu Leu Tyr Ser Asn Gly Lys Thr Tyr Leu Asn Trp Leu  55 60	
tta cag agg cca ggc cag tct cca aag cgc cta atc tat ctg gtg tct Leu Gln Arg Pro Gly Gln Ser Pro Lys Arg Leu Ile Tyr Leu Val Ser 65 70 75	242
aaa ctg gac tct gga gtc cct gac agg ttc act ggc agt gga tca gga Lys Leu Asp Ser Gly Val Pro Asp Arg Phe Thr Gly Ser Gly 80 85	290
aca gat ttt aca ctg aaa atc agc aga gtg gag gct gag gat ttg gga Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly 95	338
gtt tat tac tgc gtg caa ggt aca cat ttt cct cac acg ttc gga ggg Val Tyr Tyr Cys Val Gln Gly Thr His Phe Pro His Thr Phe Gly Gly 110 115 120	386
ggg acc aag ctg gaa ata aaa cgg gctgatgctg caccaactgt atccatcttc Gly Thr Lys Leu Glu Ile Lys Arg 125 130	440
ccaccatcca gtgagcagtt aacatctgga ggtgcctcag tcgtgtgctt cttgaacaac	500
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gtoctgaaca gttggactga toaggacago aaagacagoa cotacagoat gagcagoaco	c 680
ctcacgttga ccaaggacga gtatgaacga cataacagct atacctgtga ggccactca	729
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Thr Ile Gly Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu 35 40 45	
Leu Tyr Ser Asn Gly Lys Thr Tyr Leu Asn Trp Leu Leu Gln Arg Pro 50 55	
Gly Gln Ser Pro Lys Arg Leu Ile Tyr Leu Val Ser Lys Leu Asp Ser 65 70 75 80	
Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr 85 90 95	
Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr Tyr Cys 100 105 110	
Val Gln Gly Thr His Phe Pro His Thr Phe Gly Gly Thr Lys Leu 115 120 125	
Glu Ile Lys Arg 130	
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gag ctg gtg aag cct ggg gct tta gtg aag ata tcc tgc aag gct tct Glu Leu Val Lys Pro Gly Ala Leu Val Lys Ile Ser Cys Lys Ala Ser	145
ggt tac acc ttc aca agc tac gat ata aac tgg gtg aag cag agg cct  ggt Tyr Thr Phe Thr Ser Tyr Asp Ile Asn Trp Val Lys Gln Arg Pro  60	193
gga cag gga ctt gag tgg att gga tgg att tat cct gga gat ggt ggt Gly Gln Gly Leu Glu Trp Ile Gly Trp Ile Tyr Pro Gly Asp Gly Gly 75	241
act aat tac aat gag aaa ttc aag ggc aag gcc aca ctg act gca gac Thr Asn Tyr Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp 85	289
aaa tcc tcc agc aca gcc tac atg cag ctc agt agc ctg act tct gag  Lys Ser Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu  105	337
aac tct gca gtc tat ttc tgt gca aga ggg ggt aac ttc cct tct tat  Asn Ser Ala Val Tyr Phe Cys Ala Arg Gly Gly Asn Phe Pro Ser Tyr	385
gct atg gac tac tgg ggt caa gga acc tca gtc acc gtc tcc tca Ala Met Asp Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser 130 135	430
125 gccaaaacga caccccatc tgtctatcca ctggcccctg gatctgctgc ccaaactaac	490
gccaaaacga caccccatc tgtctatcca ctggccctts s	550
tccatggtga ccccgggatg cctggtcaag ggctatttcc ctgagccagt gacagtgacc	610
tggaactetg gatecetgte cageggtgtg cacacettee cagetgteet geagtetgae	670
ctctacactc tgagcagetc agtgactgtc cectecagca cetggeccag egagacegte	730
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gattgt	
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Pro Gly Ala Leu Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Phe 35

Thr Ser Tyr Asp Ile Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu 60 55 50

Glu Trp Ile Gly Trp Ile Tyr Pro Gly Asp Gly Gly Thr Asn Tyr Asn 75 70

Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser 85

Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu Asn Ser Ala Val 105 100

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504 <211>

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4D4 Light chain V region, plus leader <223>

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cct gct tcc aac agt gat gtt ttg atg acc caa tct cca ctc tcc ctg Pro Ala Ser Asn Ser Asp Val Leu Met Thr Gln Ser Pro Leu Ser Leu 25 30	96
cct gtc agt ctt gga gat caa gcc tcc atc tct tgc aga tct agt cag Pro Val Ser Leu Gly Asp Gln Ala Ser Ile Ser Cys Arg Ser Ser Gln 45	144
agc att gtc cat agt aat gga gac acc tat tta gaa tgg tac ctg cag Ser Ile Val His Ser Asn Gly Asp Thr Tyr Leu Glu Trp Tyr Leu Gln 50	192
aaa cca ggc cag tct cca aag ctc ctg atc tac aag gtt tcc gac cga Lys Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Lys Val Ser Asp Arg 65	240
ttt tct ggg gtc cca gac agg ttc agt ggc agt gga tca ggg aca gat Phe Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Thr Asp 80 85	288
ttc aca ctc aag atc agc aga gtg gag gct gag gat ctg gga gtt tat  Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Val Tyr  100  100	336
ttc tgc ttt caa ggt tca cat gtt ccg tac gcg ttc gga ggg ggg acc Phe Cys Phe Gln Gly Ser His Val Pro Tyr Ala Phe Gly Gly Gly Thr 125	384
aag ctg gaa ata aaa cgg gctgatgctg caccaactgt atccatcttc Lys Leu Glu Ile Lys Arg 130	432
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cct gga gcg tct ctg aaa ctc tcc tgt gca gcc tct gga ttc agt ttc Pro Gly Ala Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe 35	205
agt aac tat ggc atg tct tgg gtt cgc cag act tca gac aag agg ctg Ser Asn Tyr Gly Met Ser Trp Val Arg Gln Thr Ser Asp Lys Arg Leu 50	253
gag tgg gtc gct tcc att agt acg ggt ggt gct aat acc ttc tat cca Glu Trp Val Ala Ser Ile Ser Thr Gly Gly Ala Asn Thr Phe Tyr Pro 75	301
gac aat gta aag ggc cga ttc acc att tcc aga gag aat gcc aag aac Asp Asn Val Lys Gly Arg Phe Thr Ile Ser Arg Glu Asn Ala Lys Asn 90	349
acc cta tac ctg caa atg agt agt ctg aag tct gag gac acg gcc ttg Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Leu 100	397
tat ttc tgt gca aga gat agt cac tcc gta ggt tgt tgg ttt gct acc Tyr Phe Cys Ala Arg Asp Ser His Ser Val Gly Cys Trp Phe Ala Thr	445
tgg ggc caa ggg act ctg gtc act gtc tct gca gccaaaacaa cacccccatc Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala 135	498
agtctatcca ctggcccctg ggtgtggaga tactactggt tcctccgtga ctctgggatg	558
agtictation of geological graduation agreement agreement to cotton of the cotton of th	618
cagcagtgtg cacaccttcc cagctctcct gcagtctgga ctctacacta tgagcagctc	678
agtgactgtc ccctccagca cctggccaag ccagaccgtt acctgcagtg ttgctcaccc	738
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Pro Gly Ala Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Ser Phe

Ser Asn Tyr Gly Met Ser Trp Val Arg Gln Thr Ser Asp Lys Arg Leu 55 50

Glu Trp Val Ala Ser Ile Ser Thr Gly Gly Ala Asn Thr Phe Tyr Pro 70 65

Asp Asn Val Lys Gly Arg Phe Thr Ile Ser Arg Glu Asn Ala Lys Asn 85

Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Leu 100

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Ala Ser Glu Asn Val Val Thr Tyr Val Ser Trp Tyr Gln Gln Lys Pro 50	
Glu Gln Ser Pro Lys Leu Leu Ile Tyr Gly Ala Ser Asn Arg Tyr Thr 80 65	
Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Ala Thr Asp Phe Thr 95 85	
Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu Ala Asp Tyr His Cys 100	
Gly Gln Gly Tyr Ser Tyr Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu 115	

Glu Ile Lys Arg

•

• 130 •